=> d hist full

(FILE 'HOME' ENTERED AT 15:38:23 ON 19 APR 2006)

FILE 'REGISTRY' ENTERED AT 15:38:33 ON 19 APR 2006

L1 1 SEA ABB=ON PLU=ON 156681-44-6/RN

SET LINE 250

SET DETAIL OFF

SET NOTICE 1 DISPLAY

SET LINE LOGIN

SET DETAIL LOGIN

DIS L1 1 SQIDE

SET NOTICE LOGIN DISPLAY

FILE 'BIOSIS, BIOTECHNO, CA, CAPLUS' ENTERED AT 15:40:16 ON 19 APR 2006

FILE 'REGISTRY' ENTERED AT 15:40:24 ON 19 APR 2006

SET SMARTSELECT ON

L2 SEL PLU=ON L1 1- CHEM :

SET SMARTSELECT OFF

FILE 'BIOSIS, BIOTECHNO, CA, CAPLUS' ENTERED AT 15:40:25 ON 19 APR 2006

L3

392 SEA ABB=ON PLU=ON L2 57 SEA ABB=ON PLU=ON L3 (S) (BILE (3A) ACID OR PHYTANIC OR L4

PRISTANIC OR TRIMETHYLUNDECANOIC)

L5 27 SEA ABB=ON PLU=ON L4 AND (MEASUR? OR QUANTI? OR ACTIVIT? OR

AMOUNT)

L6 12 DUP REM L5 (15 DUPLICATES REMOVED)

D L6 IBIB ABS 1-12

FILE HOME

```
ANSWER 1 OF 1 REGISTRY COPYRIGHT 2006 ACS on STN
L1
    156681-44-6 REGISTRY
RN
CN
    Racemase, α-methylacyl coenzyme A (9CI) (CA INDEX NAME)
OTHER NAMES:
CN \alpha-Methylacyl CoA racemase
    \alpha-Methylacyl-CoA racemase
CN
CN
    2-Methylacyl-CoA racemase
MF
    Unspecified
CI
    MAN
SR
    CA
LC
    STN Files:
                 BIOSIS, CA, CAPLUS, CIN, EMBASE, TOXCENTER, USPATFULL
DT.CA Caplus document type: Conference; Dissertation; Journal; Patent
      Roles from patents: ANST (Analytical study); BIOL (Biological study);
      PREP (Preparation); PRP (Properties); USES (Uses)
      Roles from non-patents: ANST (Analytical study); BIOL (Biological
RL.NP
       study); OCCU (Occurrence); PREP (Preparation); PROC (Process); PRP
       (Properties); USES (Uses)
*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***
             99 REFERENCES IN FILE CA (1907 TO DATE)
```

100 REFERENCES IN FILE CAPLUS (1907 TO DATE)

PRINT

Mark a special word or phrase in this record:

Mark!

All organisms Homo sapiens Mus musculus

Rattus norvegicus

Mycobacterium tuberculosis

Select one or more organisms in this record:

Submit

OPCANISM COMMENTARY LITERATURE

EC NUMBER COMMNTARY

5.1.99.4

Pathway KEGG Link

No entries in this field

RECOMMENDED NAME

GeneOntology No.

alpha-Methylacyl-CoA racemase 8111

SYSTEMATIC NAME

CVNONVMC

2-Methylacyl-CoA 2-epimerase

SYNONYMS	ORGANISM	COMMENTARY	LITERATURE
2-arylpropionyl-CoA epimerase	-	-	-
2-methylacyl-CoA racemase	-	-	-
alpha-Methylacyl CoA racemase	•	-	•
GenBank U89905-derived protein GI2145184	-	-	-
GenBank U89906-derived protein GI 2145186	•	-	•
Racemase, alpha-methylacyl coenzyme A	-	-	•
Racemase, alpha-methylacyl coenzyme A (Mus musculus clone 3)	-	-	•
Racemase, alpha-methylacyl coenzyme A (Rattus norvegicus clone 11)	-		-

COMMENTARY CAS REGISTRY NUMBER

156681-44-6

197731-71-8

racemase, alpha-methylacyl coenzyme A (Mus musculus clone 3) /genBank U89906-derived

protein GI 2145186

racemase, alpha-methylacyl coenzyme A (Rattus norvegicus clone 11) /genBank U89905-derived 197731-72-9

protein GI2145184

REACTION COMMENTARY ORGANISM LITERATURE

(2S)-2-Methylacyl-CoA = (2R)-2-methylacyl-CoA

REACTION TYPE ORGANISM COMMENTARY LITERATURE

Racemization

ORGANISM COMMENTARY LITERATURE SEQUENCE CODE

 Homo sapiens
 2447, 2449

 Mus musculus
 2448

 Mycobacterium tuberculosis
 649225

 Rattus norvegicus
 2446, 2447, 2448, 2449, 649732

•					
SUBSTRATE	PRODUCT	REACTION DIAGRAM	ORGANISM	COMMENTARY/ Substrate r:=reversible ir:=irreversible	LITERATURE/ COMI Substrate Produ
(S)-2-Methylmyristoyl- CoA	(R)-2-Methylmyristoyl- CoA	<u> </u>	Rattus norvegicus	r	2446
(S)-2- Methyltetradecanoyl- CoA	(R)-2- Methyltetradecanoyl- CoA	<u> </u>	Homo sapiens	r	2 <u>447</u>
(S)-PristanoyI-CoA	(R)-PristanoyI-CoA		Homo sapiens	-	2447
(S)-PristanoyI-CoA	(R)-Pristanoyl-CoA	<u>A</u>	Rattus norvegicus	ſ	2446
(S)-PristanoyI-CoA	(R)-Pristanoyl-CoA	<u> </u>	Rattus norvegicus	r	2447
(S)- Trihydroxycoprostanoyl- CoA	(R)- Trihydroxycoprostanoyl- CoA		Homo sapiens	-	2447
(S)- Trihydroxycoprostanoyl- CoA	(R)- Trihydroxycoprostanoyl- CoA		Rattus norvegicus	•	<u>2447</u>
(S)- Trihydroxycoprostanoyl- CoA	(R)- Trihydroxycoprostanoyl- CoA	<u> </u>	Rattus norvegicus	r	<u>2446</u>
More	?		Homo sapiens	no activity with free fatty acids	<u>2447</u>
More	?		Mycobacterium tuberculosis	key enzyme in the metabolism of 2-methyl- branched fatty acids	<u>649225</u>
More	?	<u> </u>	Rattus norvegicus	•	2447
More	?		Rattus norvegicus	enzyme is involved in the alternative pathway of cholesterol sidechain oxidation. The alternative pathway consists of alphamethylacyl-CoA racemase, and peroxisomal multifunctional enzyme type 1 (peroxisomal multifunctional 2-enoyl-CoA	649732

More

hydratase/(S)-3hydroxyacyl-CoA dehydrogenase)

Rattus norvegicus no activity with 3-methylbranched or linear-chain

acyl-CoAs

NATURAL NATURAL REACTION ORGANISM COMMENTARY LITERATURE COMMENTARY LITERATURE O SUBSTRATES PRODUCTS DIAGRAM SUBSTRATE (Substrate) PRODUCT (Product) (F

No entries in this field

COFACTOR ORGANISM COMMENTARY LITERATURE IMAGE

No entries in this field

METALS and IONS ORGANISM COMMENTARY LITERATURE

No entries in this field

INHIBITORS	ORGANISM	COMMENTARY	LITERATURE	IMAGE
1-Ethyl-3-(3-dimethylaminopropyl)-carbodiimide	Rattus norvegicus	-	<u>2446</u>	● 2D- image
2-(4-lsobutylphenyl)propionic acid	Homo sapiens	-	2447	● 2D- image
2-(4-lsobutylphenyl)propionic acid	Rattus norvegicus	-	2447	● <u>2D-</u> image
2-(4-IsobutyIphenyI)propionic acid	Rattus norvegicus	i.e. ibuprofen, strongest of all competitive inhibitors tested	2446	● 2D- image
2-Methylmyristoyl-CoA	Homo sapiens	-	2447	● <u>2D-</u> image
2-Methylmyristoyl-CoA	Rattus norvegicus	-	2447	● <u>2D-</u> image
2-Methylmyristoyl-CoA	Rattus norvegicus	inhibits reaction with 2-pristanoyl-CoA	2446	● <u>2D-</u> image
2-Methyloctanoyl-CoA	Homo sapiens	•	2447	● <u>2D-</u> image
2-Methyloctanoyl-CoA	Rattus norvegicus	-	2447	● <u>2D-</u> image
2-Methyloctanoyl-CoA	Rattus norvegicus	inhibits reaction with 2-pristanoyl-CoA	2446	● 2D- image
5,5'-Dithiobis(2-nitrobenzoate)	Rattus norvegicus	inhibition is reversed by incubation of the inactivated enzyme with 10 mM dithiothreitol	2446	● 2D- image
Cu2+	Rattus norvegicus	-	2446	● 2D- image
Diethylpyrocarbonate	Rattus norvegicus	•	<u>2446</u>	● <u>2D-</u> image
Diisopropylphosphofluoridate	Rattus norvegicus	•	2446	● <u>2D-</u> image
Fe2+	Rattus norvegicus	slight inhibition	2446	● 2D- image
Hg2+	Rattus	-	2446	● <u>2D-</u>

	no	orvegicus					<u>image</u>
More		attus orvegicus	no inhi	bition by iodoace	tamide	2446	-
NEM		attus orvegicus	weak			2446	● <u>2D-</u> image
Palmitoyl-CoA	Ho			on is caused by t mixed micelles	he formation of	2447	● 2D- image
PalmitoyI-CoA		attus orvegicus		-		2447	● 2D- image
Palmitoyl-CoA				ites at low conce 0.1 mM	ntrations, inhibits	<u>2446</u>	● 2D- image
thimerosal		attus orvegicus	slight			2446	● 2D- image
ACTIVATING COMPOUND	ORGANISM	COMMEN	TARY		1	LITERATUR	E IMAGE
Palmitoyl-CoA	Rattus norvegicus	stimulates 0.1 mM	at low	concentrations, i	nhibits above	<u>2446</u>	● <u>2D-</u> image
KM VALUE KM VALUE [mM] Maximum	E [mM] SUBS	STRATE		ORGANISM	COMMENTARY	LITERATU	RE IMAGE
0.076 -	pristar	noyl-CoA		Rattus norvegicus	-	<u>2446</u>	● 2D- image
0.172 -	pristar	noyl-CoA		Homo sapiens	-	<u>2447</u>	● <u>2D-</u> image
0.0316 -	Trihyd CoA	droxycoprostar	noyl-	Homo sapiens	-	<u>2447</u>	● <u>2D-</u> image
0.06 -	Trihyd CoA	droxycoprostar	noyl-	Rattus norvegicus	-	2446	● <u>2D-</u> image
Ki VALUE [mM] Ki VAL	UE (mM) Maximi	um INHIBITO	R OR	GANISM COMM	MENTARY LITER	ATURE IM	AGE
No entries in this field							
	URNOVER NUM IAXIMUM[1/s]	BER S	SUBST	RATE ORGANI	SM COMMENTA	RY LITERA	TURE IMAGE
No entries in this field							
ACTIVITY ACT	CIFIC IVITY IMUM	ORGANISM	CON	MMENTARY			LITERATURE
additional information	-	Homo sapien			upled assay based idase/peroxidase	d on the use	2449
additional information	-	Rattus norvegicus			upled assay based idase/peroxidase	d on the use	2449
additional information		Rattus norvegicus		elopment of a ve ometric assay	ry sensitive and co	onvenient	2446
pH OPTIMUM pH MAXII	MUM ORGANIS	м сом	MENT	ARY	LITERATURE		
8 -	Homo sapi	iens		•	2447		
7 -	Rattus nor	vegicus trihyd	droxyco	prostanoyl-CoA	2446		
6 -	Rattus nor	vegicus prista	anoyl-C	οΑ	2446		

pH RANGE	pH RANGE MAXIMUM	ORGANISM	COMMENTARY	LITERATURE
6.5	9	Homo sapiens	more than 80% of maximal activity between pH 6.5 and pH 9.0, inactive below pH 5	2447

TEMPERATURE OPTIMUM TEMPERATURE OPTIMUM MAXIMUM ORGANISM COMMENTARY LITERATURE No entries in this field

TEMPERATURE RANGE TEMPERATURE MAXIMUM ORGANISM COMMENTARY LITERATURE No entries in this field

SOURCE TISSUE	ORGANISM	COMM	ENTARY	LITERATURE
adrenal gland	Homo sapiens	weak		2449
cerebellum	Homo sapiens	weak		2449
chorionic villus	Homo sapiens		-	2447
fibroblast	Homo sapiens		-	2447
harderian gland	Homo sapiens		-	2449
heart	Homo sapiens	weak		2449
Hep-G2 cell	Homo sapiens		-	2447
intestinal mucosa	Homo sapiens		-	2449
kidney	Homo sapiens		•	2449
liver	Homo sapiens		-	<u>2447, 2449</u>
liver	Rattus norvegicus		-	<u>2446, 2447, 2449</u>
lung .	Homo sapiens	weak		2449
muscle	Homo sapiens	weak		2449
pancreas	Homo sapiens	weak		2449
skin fibroblast	Homo sapiens		-	2447
spleen	Homo sapiens	weak		2449
telencephalon	Homo sapiens	weak		2449
testis	Homo sapiens	weak		2449
thymus	Homo sapiens	weak		2449

LOCALIZATION	ORGANISM	COMMENTARY			GeneO No.	ntology	LITERATURE
mitochondrion	Homo sapiens	only 10-30% of the activity is fo	und in m	nitochondria	5739		<u>2447</u>
mitochondrion	Rattus norvegicus	co-distributed exclusively with a enzymes	mitochon	drial marker	5739		2447
peroxisome	Rattus norvegicus	-			<u>5777</u>		649732
peroxisome	Rattus norvegicus	bulk activity			<u>5777</u>		2447
ACCESSION CODE	ENTRY NAME	ORGANISM	NO. OF AA	MOLECULA WEIGHT[Da		SOURCE	Sequence
Q9UHK6 pBLAST	AMACR_HUMAN	Homo sapiens	382	42360		Swiss- Prot	<u>Show</u> Sequence

O09174 pBLAST	AMACR_MOUSE	Mus musculus	380	41587	Swiss- Prot	Show Sequence
P70473 pBLAST	AMACR_RAT	Rattus norvegicus	381	41697	Swiss- Prot	U Show Sequence
Q4IYP2 pBLAST	Q4IYP2_AZOVI	Azotobacter vinelandii AvOP	397	42454	TrEMBL	Show Sequence
Q4LND9 pBLAST	Q4LND9_9BURK	Burkholderia cenocepacia HI2424	455	49064	TrEMBL	U Show Sequence
Q4LQP6 pBLAST	Q4LQP6_9BURK	Burkholderia cenocepacia HI2424	350	36615	TrEMBL	<u> </u>
Q4LWN5 pBLAST	Q4LWN5_9BURK	Burkholderia cenocepacia HI2424	406	43669	TrEMBL	Show Sequence
Q4LXS2 pBLAST	Q4LXS2_9BURK	Burkholderia cenocepacia HI2424	406	43786	TrEMBL	<u> Show</u> Sequence
Q4NIB3 pBLAST	Q4NIB3_9MICC	Arthrobacter sp. FB24	419	45712	TrEMBL	₫ <u>Show</u> Sequence
Q4NR64 pBLAST	Q4NR64_9DELT	Anaeromyxobacter dehalogenans 2CP-C	391	40575	TrEMBL	⊈ <u>Show</u> Sequence
Q4AQK8 pBLAST	Q4AQK8_9BURK	Polaromonas sp. JS666	433	45750	TrEMBL	<u> Show</u> Sequence
Q4ASU3 pBLAST	Q4ASU3_9BURK	Polaromonas sp. JS666	407	43920	TrEMBL	<u> </u>
Q4AVI5 pBLAST	Q4AVI5_9BURK	Polaromonas sp. JS666	416	45848	TrEMBL	<u>[♣] Show</u> Sequence
Q4B2Y0 pBLAST	Q4B2Y0_9BURK	Polaromonas sp. JS666	407	43807	TrEMBL	Ů <u>Show</u> Sequence
Q4B520 pBLAST	Q4B520_9BURK	Polaromonas sp. JS666	416	44626	TrEMBL	Show Sequence
Q447U6 pBLAST	Q447U6_SOLUS	Solibacter usitatus Ellin6076	403	44029	TrEMBL	Show Sequence
Q422R9 pBLAST	Q422R9_DESHA	Desulfitobacterium hafniense DCB-2	355	39734	TrEMBL	<u>[₾] Show</u> Sequence
Q3VZH3 pBLAST	Q3VZH3_9ACTO	Frankia sp. EAN1pec	423	44956	TrEMBL	Show Sequence
Q3VZL8 pBLAST	Q3VZL8_9ACTO	Frankia sp. EAN1pec	379	40254	TrEMBL	Show Sequence
Q3W154 pBLAST	Q3W154_9ACTO	Frankia sp. EAN1pec	378	40088	TrEMBL	<u>[♣] Show</u> Sequence
Q3W4K6 pBLAST	Q3W4K6_9ACTO	Frankia sp. EAN1pec	369	39541	TrEMBL	Show Sequence
Q3W562 pBLAST	Q3W562_9ACTO	Frankia sp. EAN1pec	396	43121	TrEMBL	Show Sequence
Q3W5F2 pBLAST	Q3W5F2_9ACTO	Frankia sp. EAN1pec	451	48468	TrEMBL	Ú Show Sequence
Q3WFQ7 pBLAST	Q3WFQ7_9ACTO	Frankia sp. EAN1pec	402	43090	TrEMBL	Ů Show Sequence
Q3WFX1 pBLAST	Q3WFX1_9ACTO	Frankia sp. EAN1pec	389	41842	TrEMBL	Show Sequence
Q3WHJ3 pBLAST	Q3WHJ3_9ACTO	Frankia sp. EAN1pec	462	49208	TrEMBL	Ú <u>Show</u> Sequence
Q3WJ86 pBLAST	Q3WJ86_9ACTO	Frankia sp. EAN1pec	771	81365	TrEMBL	Show Sequence
Q3WTP8	Q3WTP8_9RHIZ	Mesorhizobium sp. BNC1	419	45059	TrEMBL	⊕ Show

pBLAST						Sequence
Q3WTR2 pBLAST	Q3WTR2_9RHIZ	Mesorhizobium sp. BNC1	364	38783	TrEMBL	Show Sequence
Q3WXX8 pBLAST	Q3WXX8_9RHIZ	Mesorhizobium sp. BNC1	378	40584	TrEMBL	<u>Show</u> Sequence
Q3X096 pBLAST	Q3X096_9ACTN	Rubrobacter xylanophilus DSM 9941	404	44338	TrEMBL	<u> Show</u> Sequence
Q3X5E1 pBLAST	Q3X5E1_9ACTN	Rubrobacter xylanophilus DSM 9941	395	43719	TrEMBL	Ů <u>Show</u> Sequence
Q3X5K5 pBLAST	Q3X5K5_9ACTN	Rubrobacter xylanophilus DSM 9941	414	45231	TrEMBL	<u> </u>
Q3MVD3 pBLAST	Q3MVD3_9DELT	Syntrophobacter fumaroxidans MPOB	391	42948	TrEMBL	Ů <u>Show</u> Sequence
Q3QQ94 pBLAST	Q3QQ94_9RHOB	Silicibacter sp. TM1040	373	40180	TrEMBL	U Show Sequence
Q3QTP0 pBLAST	, Q3QTP0_9RHOB	Silicibacter sp. TM1040	395	42380	TrEMBL	<u> Show</u> Sequence
Q3FMR0 pBLAST	Q3FMR0_9BURK	Rhodoferax ferrireducens DSM 15236	362	38569	TrEMBL	<u>[©] Show</u> Sequence
Q3FP81 pBLAST	Q3FP81_9BURK	Rhodoferax ferrireducens DSM 15236	418	44695	TrEMBL	<u> Show</u> Sequence
Q3FU66 pBLAST	Q3FU66_9BURK	Rhodoferax ferrireducens DSM 15236	387	41780	TrEMBL	<u> Show</u> Sequence
Q3GL65 pBLAST	Q3GL65_9GAMM	Psychrobacter cryohalolentis K5	352	38399	TrEMBL	<u> Show</u> Sequence
Q3GNY3 pBLAST	Q3GNY3_9GAMM	Psychrobacter cryohalolentis K5	423	45301	TrEMBL	Show Sequence
Q3K8D2 pBLAST	Q3K8D2_PSEFL	Pseudomonas fluorescens PfO-1	393	42243	TrEMBL	Ú <u>Show</u> Sequence
Q391W0 pBLAST	Q391W0_9BURK	Burkholderia sp. 383	388	41422	TrEMBL	<u>[©] Show</u> Sequence
Q392M2 pBLAST	Q392M2_9BURK	Burkholderia sp. 383	381	40921	TrEMBL	Ů <u>Show</u> Sequence
Q39B91 pBLAST	Q39B91_9BURK	Burkholderia sp. 383	350	36647	TrEMBL	<u> </u>
Q39DK7 pBLAST	Q39DK7_9BURK	Burkholderia sp. 383	463	49229	TrEMBL	<u>⁴ Show</u> Sequence
Q39LX9 pBLAST	Q39LX9_9BURK	Burkholderia sp. 383	384	40897	TrEMBL	<u> </u>
Q39MI0 pBLAST	Q39MI0_9BURK	Burkholderia sp. 383	388	41225	TrEMBL	Show Sequence
Q39MT5 pBLAST	Q39MT5_9BURK	Burkholderia sp. 383	350	37383	TrEMBL	Show Sequence
Q39MW4 pBLAST	Q39MW4_9BURK	Burkholderia sp. 383	386	41428	TrEMBL	Show Sequence
Q39N02 pBLAST	Q39N02_9BURK	Burkholderia sp. 383	369	38590	TrEMBL	Ú <u>Show</u> Sequence
Q3CNG3 pBLAST	Q3CNG3_ALTAT	Pseudoalteromonas atlantica T6c	386	42619	TrEMBL	Show Sequence
Q8F1J1 pBLAST	Q8F1J1_LEPIN	Leptospira interrogans	390	43625	TrEMBL	Show Sequence
Q89XH6 pBLAST	Q89XH6_BRAJA	Bradyrhizobium japonicum	388	41545	TrEMBL	Show Sequence

O06543 pBLAST	O06543_MYCTU	Mycobacterium tuberculosis	360	38685	TrEMBL	Ú <u>Show</u> Sequence
Q6FBN2 pBLAST	Q6FBN2_ACIAD	Acinetobacter sp. (strain ADP1)	407	45097	TrEMBL	Show Sequence
Q6FBN5 pBLAST	Q6FBN5_ACIAD	Acinetobacter sp. (strain ADP1)	405	44587	TrEMBL	Ú Show Sequence
Q7U0J6 pBLAST	Q7U0J6_MYCBO	Mycobacterium bovis	360	38685	TrEMBL	Show Sequence
Q8YB25 pBLAST	Q8YB25_BRUME	Brucella melitensis	405	43553	TrEMBL	Show Sequence
Q8YB81 pBLAST	Q8YB81_BRUME	Brucella melitensis	415	45073	TrEMBL	Show Sequence

PDB

ORGANISM

1X74, download Mycobacterium tuberculosis

MOLECULAR WEIGHT	MOLECULR WEIGHT MAXIMUM	ORGANISM	COMMENTARY	LITERATURE
75000	-	Mycobacterium tuberculosis	dynamic light-scattering measurement	649225
47700	-	Homo sapiens	gel filtration	<u>2447</u>
44700		Rattus norvegicus	gel filtration	2446

SUBUNITS ORGANISM

COMMENTARY

LITERATURE

Dimer

Mycobacterium tuberculosis 2 * 39000, SDS-PAGE 649225

Monomer Homo sapiens

1 * 47100, SDS-PAGE 2447

Monomer Rattus norvegicus

2447

Monomer Rattus norvegicus

1 * 44900, SDS-PAGE 2446

POSTTRANSLATIONAL MODIFICATION ORGANISM COMMENTARY LITERATURE

No entries in this field

Crystallization/COMMENTARY

ORGANISM

LITERATURE

hanging-drop vapour-diffusion method, the best crystals grow in 1.26 M ammonium phosphate, pH 7.0 using a protein concentration of 24 mg/ml

Mycobacterium

649225

tuberculosis

PH STABILITY PH STABILITY MAXIMUM ORGANISM COMMENTARY LITERATURE

No entries in this field

TEMPERATURE STABILITY TEMPERATURE STABILITY MAXIMUM ORGANISM

COMMENTARY

LITERATURE

50 35

40

Homo sapiens half-life: 15 min

2447

Homo sapiens slow loss of activity 2447

GENERAL STABILITY ORGANISM LITERATURE

No entries in this field

ORGANIC SOLVENT ORGANISM COMMENTARY LITERATURE

No entries in this field

2447

OXIDATION STABILITY ORGANISM LITERATURE

No entries in this field

STORAGE STABILITY ORGANISM LITERATURE

No entries in this field

Purification/COMMENTARY ORGANISM LITERATURE

Homo sapiens 2447

- Rattus norvegicus 2446, 2448

Mycobacterium tuberculosis <u>649225</u>

Cloned/COMMENTARY ORGANISM LITERATURE

Mus musculus 2448

expression in Escherichia coli Rattus norvegicus 2448

ENGINEERING ORGANISM COMMENTARY LITERATURE

No entries in this field

Renatured/COMMENTARY ORGANISM LITERATURE

No entries in this field

APPLICATION ORGANISM COMMENTARY LITERATURE

medicine Homo the activity of EC 5.1.99.4 may prove to be a valuable parameter for the

sapiens prenatal diagnosis of general defects of peroxisome biogenesis such as

Zellweger syndrome

DISEASE TITLE OF PUBLICATION LINK TO PUBMED

No entries in this field

REF.	AUTHORS	TITLE	JOURNAL	VOL.	PAGES	YEAR	ORGANISM	COMMENTARY	LINK TO PUBMED
2446	Fingerhut, R.;	Purification and properties of an alpha-methylacyl- CoA racemase from rat liver	Eur. J. Biochem.	222	313- 323	1994	Rattus norvegicus	c	● <u>PubM</u> €
2447	Schmitz, W.; Albers, C.; Fingerhut, R.; Conzelmann, E.	Purification and characterization of an alpha-methylacyl-CoA racemase from human liver	Eur. J. Biochem.	231	815- 822	1995	Homo sapiens, Rattus norvegicus	С	● <u>PubM</u> €
2448	Schmitz, W.; Helander, H.M.; Hiltunen, J.K.; Conzelmann, E.	Molecular cloning of cDNA species for rat and mouse liver alpha- methylacyl-CoA racemase	Biochem. J.	326	883- 889	1997	Mus musculus, Rattus norvegicus	С	● <u>PubMe</u>
2449	Van	2-Methylacyl	Biochim.	1347	62-68	1997	Homo sapiens,	c .	● PubMe

	Veldhoven, P.P.; Croes, K.; Casteels, M.; Mannaerts, G.P.	racemase: a coupled assay based on the use of pristanoyl-CoA oxidase/peroxidase and reinvestigation of its subcellular distribution in rat and human liver	Biophys. Acta				Rattus norvegicus		
649225	Bhaumik, P.; Kursula, P.; Ratas, V.; Conzelmann, E.; Hiltunen, J.K.; Schmitz, W.; Wierenga, R.K.	Crystallization and preliminary X-ray diffraction studies of an alpha- methylacyl-CoA racemase from Mycobacterium tuberculosis	Acta Crystallogr. Sect. D	59	353- 355	2003	Mycobacterium tuberculosis	C	● <u>PubM</u> e
649732	Cuebas, D.A.; Phillips, C.; Schmitz, W.; Conzelmann, E.; Novikov, D.K.	The role of alpha- methylacyl-CoA racemase in bile acid synthesis	Biochem. J.	363	801- 807	2002	Rattus norvegicus	С	● <u>PubMe</u>

LINKS TO OTHER DATABASES (specific for EC-Number 5.1.99.4)

ExPASy

KEGG

NCBI: PubMed, Protein, Nucleotide, Structure, Genome, OMIM, Domains

IUBMB Enzyme Nomenclature

PDB database(3D structure)

PROSITE Database of protein families and domains

SYSTERS

Protein Mutant Database

Structural Classification of Proteins (SCOP)

Protein Structure Classification (CATH)

InterPro (database of protein families, domains and functional sites)